

AMENDMENTS

Amendments to the Claims:

Listing of Claims:

The following listing of claims replaces all previous listings or versions thereof:

1 - 18. (canceled)

19. (previously presented) A method of producing a virus comprising:

introducing into a host cell a recombinant GBV-B or chimeric GBV-B viral genome comprising a 3' terminal sequence of GBV-B, wherein the 3' terminal sequence comprises 50 contiguous nucleotides from SEQ ID NO:1; and culturing said host cell under conditions permitting production of a virus from said genome.

20. (previously presented) The method of claim 19, wherein said 3' terminal sequence comprises 100 contiguous nucleotides from SEQ ID NO:1.

21. (previously presented) The method of claim 20, wherein said 3' terminal sequence comprises SEQ ID NO:1.

22 - 28. (canceled)

29. (original) The method of claim 28, wherein said host cell is in an animal.

30. (previously presented) The method of claim 19, wherein said genome comprises recombinant RNA.

31. (previously presented) The method of claim 19, wherein said genome is encoded by recombinant DNA.

32. (original) The method of claim 19, further comprising the step of isolating virus from said host cell.

33. (original) The method of claim 32, wherein said virus is purified to homogeneity.

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34 - 50. (canceled)

51. - 55. (canceled)

56. (previously presented) A method of producing a GBV-B or chimeric GBV-B virus comprising:

obtaining a virus produced by the method of claim 19;

introducing the virus into a second host cell; and

culturing said host cell under conditions permitting production of virus.

57. (previously presented) A method of producing a virus comprising:

introducing into a host cell a recombinant GBV-B or chimeric GBV-B viral genome comprising a 3' terminal sequence of GBV-B, wherein the 3' terminal sequence is at least 70% identical to SEQ ID NO:1; and

maintaining said host cell under conditions permitting production of a virus from said genome.

58. (previously presented) The method of claim 57, wherein the host cell is a liver cell.

59. (previously presented) The method of claim 58, wherein the liver cell is in an animal.

60. (previously presented) The method of claim 59, wherein the animal is a primate.

61. (previously presented) The method of claim 60, wherein the primate is a non-human primate.

62. (previously presented) The method of claim 60, wherein the primate is a tamarin.

63. (previously presented) The method of claim 58, wherein the 3' terminal sequence is at least 80% identical to SEQ ID NO:1.

64. (previously presented) The method of claim 63, wherein the 3' terminal sequence is at least 90% identical to SEQ ID NO:1.

65. (previously presented) The method of claim 64, wherein the 3' terminal sequence is at least 95% identical to SEQ ID NO:1.
66. (previously presented) The method of claim 65, wherein the 3' terminal sequence is SEQ ID NO:1.
67. (previously presented) A method of producing a virus comprising:
introducing into a liver cell a recombinant GBV-B or chimeric GBV-B viral genome comprising a 3' terminal sequence of GBV-B, wherein the 3' terminal sequence is at least 70% identical to SEQ ID NO:1; and
maintaining said liver cell under conditions permitting production of a virus from said genome.
68. (previously presented) A method of replicating a virus comprising:
introducing into a liver cell a recombinant GBV-B or chimeric GBV-B viral genome comprising a 3' terminal sequence of GBV-B, wherein the 3' terminal sequence is at least 70% identical to SEQ ID NO:1; and
maintaining said liver cell under conditions permitting replication of the recombinant viral genome.
69. (previously presented) The method of claim 68, wherein the 3' terminal sequence is at least 80% identical to SEQ ID NO:1.
70. (previously presented) The method of claim 69, wherein the 3' terminal sequence is at least 90% identical to SEQ ID NO:1.
71. (previously presented) The method of claim 70, wherein the 3' terminal sequence is at least 95% identical to SEQ ID NO:1.
72. (previously presented) The method of claim 71, wherein the 3' terminal sequence is SEQ ID NO:1.